SEQUENCE LISTING

<110> DeJong Dr., Jeff <120> Transcription Factors Related to TFIIA <130> 119941-1053 <140> <141> <160> 19 <170> PatentIn Ver. 2.0 <210> 1 <211> 1617 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (16)..(1449) <400> 1 gctggaggtg ctgtc atg gcc tgc ctc aac ccg gtg cct aaa ctc tac aga 51 Met Ala Cys Leu Asn Pro Val Pro Lys Leu Tyr Arg 1 10 tot gta att gaa gat gta att gaa gga gtt cgg aat cta ttt gct gaa 99

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Ile	Gln	Ser	Pro	Leu	Phe	Thr	Leu	Gln	Leu	Pro	His	Ser	Leu	His	Gln	
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Thr	Leu	Gln	Ser	Ser	Thr	Ala	Ser	Leu	Val	Ile	Pro	Ala	Gly	Arg	Thr	
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ctt	cca	agt	ttt	acc	aca	gca	gaa	ctg	ggc	act	tca	aac	tcc	agt	gca	339
Leu	Pro	Ser	Phe	Thr	Thr	Ala	Glu	Leu	Gly	Thr	Ser	Asn	Ser	Ser	Ala	
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aac	ttt	act	ttt	cct	ggt	tat	ссс	att	cat	gta	сса	gca	ggt	gtg	aca	387
Asn	Phe	Thr	Phe	Pro	Gly	Tyr	Pro	Ile	His	Val	Pro	Ala	Gly	Val	Thr	
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cta	cag	act	gta	tct	ggt	cac	ctt	tat	aaa	gtc	aat	gta	сса	att	atg	435
Leu	Gln	Thr	Val	Ser	Gly	His	Leu	Tyr	Lys	Val	Asn	Val	Pro	Ile	Met	
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gtg aca gag act tct gga aga gca ggt att ctt cag cat cca att cag 483
Val Thr Glu Thr Ser Gly Arg Ala Gly Ile Leu Gln His Pro Ile Gln

145 150 155

with the contrast of the section (x,y) , which is the contrast of the section (x,y)

cca caa ttg aat cca tgg tct ctt caa gca act act gaa aaa tca cag Pro Gln Leu Asn Pro Trp Ser Leu Gln Ala Thr Thr Glu Lys Ser Gln aga att gaa acc gtg cta cag caa ccc gca att cta cct tct ggg cca Arg Ile Glu Thr Val Leu Gln Gln Pro Ala Ile Leu Pro Ser Gly Pro gta gat agg aaa cac tta gaa aat gcc acc agt gat ata ctt gta tct Val Asp Arg Lys His Leu Glu Asn Ala Thr Ser Asp Ile Leu Val Ser cct gga aat gag cat aaa atc gtg cct gaa gct ttg ttg tgt cat cag Pro Gly Asn Glu His Lys Ile Val Pro Glu Ala Leu Leu Cys His Gln gaa agt tot cac tat atc agt ott coa ggt gtt gta ttt tot coa cag Glu Ser Ser His Tyr Ile Ser Leu Pro Gly Val Val Phe Ser Pro Gln gtc tct caa aca aat tct gat gtg gag tca gtg ctc agt ggt tca gct Val Ser Gln Thr Asn Ser Asp Val Glu Ser Val Leu Ser Gly Ser Ala age atg get caa aat etg cat gat gag tee etc tee aca age eet eat Ser Met Ala Gln Asn Leu His Asp Glu Ser Leu Ser Thr Ser Pro His ggg gct ctc cac cag cac gtg act gat att cag ctt cat att ctt aaa

aat agg atg tat gga tgt qat tot gta aag caa ooa aga aat ata qa $oldsymbol{q}=463$

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Glu	Asp	Pro	Leu	Asn	Ser	Gly	Asp	Asp	Val	Ser	Glu	Gln	Asp	Val	Pro	
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Asp Lou Fig. Asp Thr Asp Ash (a. 116) at (), our Tyr Asp (), (116)

 cat
 cga
 agc
 aag
 aaa
 tgg
 aaa
 ttg
 aaa
 gat
 tggt
 gtt
 atg
 1395

 His
 Arg
 Ser
 Lys
 Asn
 Lys
 Phe
 Tyr
 Leu
 Lys
 Asp
 Gly
 Val
 Met

 445
 455
 455
 460

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Glu Gln Val Leu Lys Asp Leu Lys Gln Leu Trp Glu Thr Lys Val Leu

the output of the state of the contract of the

Leu Phe Thr Leu Gln Leu Pro His Ser Leu His Gln Thr Leu Gln Ser

65 70 75 80

Ser Thr Ala Ser Leu Val Ile Pro Ala Gly Arg Thr Leu Pro Ser Phe
85 90 95

Thr Thr Ala Glu Leu Gly Thr Ser Asn Ser Ser Ala Asn Phe Thr Phe
100 105 110

Pro Gly Tyr Pro Ile His Val Pro Ala Gly Val Thr Leu Gln Thr Val

Ser Gly His Leu Tyr Lys Val Asn Val Pro Ile Met Val Thr Glu Thr

Ser Gly Arg Ala Gly Ile Leu Gln His Pro Ile Gln Gln Val Phe Gln
145 150 155 160

Gln Leu Gly Gln Pro Ser Val Ile Gln Thr Ser Val Pro Gln Leu Asn 165 170 175

Pro Trp Ser Leu Gln Ala Thr Thr Glu Lys Ser Gln Arg Ile Glu Thr
180 185 190

Val Leu Gln Gln Pro Ala Ile Leu Pro Ser Gly Pro Val Asp Arg Lys
195 200 205

His Leu Glu Asn Ala Thr Ser Asp Ile Leu Val Ser Pro Gly Asn Glu 210 215 220

290 295 300

Gly Cys Asp Ser Val Lys Gln Pro Arg Asn Ile Glu Glu Pro Ser Asn 305 310 315 320

Ile Pro Val Ser Glu Lys Asp Ser Asn Ser Gln Val Asp Leu Ser Ile
325 330 335

Arg Val Thr Asp Asp Asp Ile Gly Glu Ile Ile Gln Val Asp Gly Ser 340 345 350

Gly Asp Thr Ser Ser Asn Glu Glu Ile Gly Ser Thr Arg Asp Ala Asp
355 360 365

Glu Asn Glu Phe Leu Gly Asn Ile Asp Gly Gly Asp Leu Lys Val Pro 370 375 380

Glu Glu Glu Ala Asp Ser Ile Ser Asn Glu Asp Ser Ala Thr Asn Ser 385 390 395 400

420 425 430

Thr Asp Asn Val Ile Val Cys Gln Tyr Asp Lys Ile His Arg Ser Lys 435 440 445

Asn Lys Trp Lys Phe Tyr Leu Lys Asp Gly Val Met Cys Phe Gly Gly

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Met 1

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Val Gln Ser Ser Gln Lys Ser Lys Asn Phe Fro Leu Glu Asn Gln Gly

20 25 30

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Ile	Val	Asp	Phe	Tyr	Phe	Ser	Pro	Gly	Pro	Pro	Ser	Asn	Ser	Pro	Leu	
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Gly	Thr	His	Val	Leu	Tyr	Pro	Ile	Pro	Glu	Ser	Ser	Ser	Asp	Ser	Pro	
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Leu	Ala	Ile	Ser	Gly	Gly	Glu	Ser	Ser	Leu	Leu	Pro	Thr	Arg	Pro	Thr	
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Glu	Asp	Cys	Ala	Phe	Ser	Ser	Pro	Phe	Arg	ГЛЗ	Asp	Glu	Gly	Ser	Asp	
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Gln	Asp	Ser	Leu	Arg	Ser	Leu	Ser	Met	His	Cys	Leu	Cys	Ala	Glu	Glu	
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Asn	Ala		Ser	Phe	Val	Pro		Thr	Leu	Phe	Arg	,	Gln	Pro	Lys	
		260					265					270				
.																
			tct											_	-	981
ser	сту	Trp	Ser	rne	Met	геп	Arg	He	Pro	Glu	Lys	гЛs	Asn	Met	Met	

Ser Ser Arg Gln Trp Gly Pro Ile Phe Leu Lys Val Leu Pro Gly Gly

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ata	cag	ctt	gat	сса	tat	tgt	agg	ctt	tct	gaa	ccc	aag	gtt	gag	aac	1125
Ile	Gln	Leu	Asp	Pro	Tyr	Cys	Arg	Leu	Ser	Glu	Pro	Lys	Val	Glu	Asn	
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ttc	agt	gta	gca	gga	aaa	atc	cac	act	gtg	aag	att	gaa	cat	gtg	tct	1173
Phe	Ser	Val	Ala	Gly	Lys	Ile	His	Thr	Val	Lys	Ile	Glu	His	Val	Ser	
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Туr	Thr	Glu	Lys	Arg	Lys	Tyr	His	Ser	Lys	Thr	Glu	Val	Val	His	Glu	
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Pro	Asp	Ile	Glu	Gln	Met	Leu	Lys	Leu	Gly	Ser	Thr	Ser	Tyr	His	Asp	
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ttc	ctt	gac	ttt	ctg	act	act	gtg	gag	gag	gag	ctg	atg	aag	ttg	cca	1317
Phe	Leu	Asp	Phe	Leu	Thr	Thr	Val	Glu	Glu	Glu	Leu	Met	Lys	Leu	Pro	
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gct	gtt	tca	aaa	cca	aaa	aag	aac	tac	gag	gag	caa	gaa	att	tcc	ttg	1365
Ala	Val	Ser	Lys	Pro	Lys	Lys	Asn	Tyr	Glu	Glu	Gln	Glu	Ile	Ser	Leu	

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Val	Glu	Ser	Ala	Val	Ile	Thr	Gln	Ile	Tyr	Cys	Leu	Суз	Phe	Val	Asn	
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ggg	aac	ctg	gaa	tgc	ttt	tta	acc	ttg	aat	gac	ctt	gag	ttg	ccg	aag	1509
Gly	Asn	Leu	Glu	Суз	Phe	Leu	Thr	Leu	Asn	Asp	Leu	Glu	Leu	Pro	Lys	
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cga	gat	gaa	tcc	tat	tat	gag	aag	gac	tca	gaa	aaa	aag	ggg	att	gat	1557
Arg	Asp	Glu	Ser	Tyr	Tyr	Glu	Lys	Asp	Ser	Glu	Lys	Lys	Gly	Ile	Asp	
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Ile	Leu	Asp	Tyr	His	Phe	His	Lys	Cys	Val	Asn	Val	Gln	Glu	Phe	Glu	
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Gln	Ser	Arg	Ile	Ile	Lys	Phe	Val	Pro	Leu	Asp	Ala	Cys	Arg	Phe	Glu	
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Leu	Met	Arg	Phe	Lys	Thr	Leu	Tyr	Asn	Gly	Asp	Asn	Leu	Pro	Phe	Ser	
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Leu	Lys	Ser	Val	Val	Val	Val	Gln	Gly	Ala	Tyr	Val	Glu	Leu	Gln	Ala	
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Phe	Val	Asn	Met		Ser	Leu	Ala	Gln	Arg	Ser	Ser	Tyr	Ala	Gly	Ser	

Leu Arg Ser Sys Asp Asn Ile Arg Ile His Phe Pro Val Pro Ser Gln

565	570	575

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Lys	Ala	Lys	Met	Asn	Arg	Arg	Ala	Суз	Leu	Gly	Ser	Leu	Gln	Glu	Leu	
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Asn	Ser	Ser	Leu	Asp	His	Pro	His	Cys	Leu	Ser	Tyr	Lys	Leu	Glu	Leu	
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gga	tca	gac	caa	gaa	att	ссс	tct	gat	tgg	tat	сса	ttt	gct	act	gtt	2133
Gly	Ser	Asp	Gln	Glu	Ile	Pro	Ser	Asp	Trp	Tyr	Pro	Phe	Ala	Thr	Val	
		660					665					670				
cag	ttt	tcc	gtg	cct	gac	acc	tgt	gcc	tca	agg	aca	gag	gtc	agg	tct	2181
Gln	Phe	Ser	Val	Pro	Asp	Thr	Суѕ	Ala	Ser	Arg	Thr	Glu	Val	Arg	Ser	
	675					680					685					

gct tgc tac aac atc cag cct aaa ctc tac aga tct gta att gaa gat Ala Cys Tyr Asn Ile Gln Pro Lys Leu Tyr Arg Ser Val Ile Glu Asp gta att gaa gga gtt cgg aat cta ttt gct gaa gaa ggt ata gag gaa Val Ile Glu Gly Val Arg Asn Leu Phe Ala Glu Glu Gly Ile Glu Glu caa gtt tta aaa gac ttg aag cag ctc tgg gaa acc aag gtt ttg cag Gln Val Leu Lys Asp Leu Lys Gln Leu Trp Glu Thr Lys Val Leu Gln tct aaa gca aca gaa gac ttc ttc aga aat agc atc caa tca cct ctg Ser Lys Ala Thr Glu Asp Phe Phe Arg Asn Ser Ile Gln Ser Pro Leu ttt act ctt cag ttg ccg cac agc ttg cac caa aca ttg caa tcg tca Phe Thr Leu Gln Leu Pro His Ser Leu His Gln Thr Leu Gln Ser Ser aca gca tca tta gtt att cct gct ggt aga act ctt cca agt ttt acc Thr Ala Ser Leu Val Ile Pro Ala Gly Arg Thr Leu Pro Ser Phe Thr aca gca gaa ctg ggc act tca aac tcc agt gca aac ttt act ttt cct Thr Ala Glu Leu Gly Thr Ser Asn Ser Ser Ala Asn Phe Thr Phe Pro ggt tat ccc att cat gta cca gca ggt gtg aca cta cag act gta tct Gly Tyr Pro Ile His Val Pro Ala Gly Val Thr Leu Gln Thr Val Ser

Gly His Leu Tyr Lys Val Asn Val Pro Ile Met Val Thr Gla Thr Ger

835	840	845

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Trp	Ser	Leu	Gln	Ala	Thr	Thr	Glu	Lys	Ser	Gln	Arg	Ile	Glu	Thr	Val	
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cta	cag	caa	ссс	gca	att	cta	cct	tct	ggg	сса	gta	gat	agg	aaa	cac	2853
Leu	Gln	Gln	Pro	Ala	Ile	Leu	Pro	Ser	Gly	Pro	Val	Asp	Arg	Lys	His	
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		aat			_	gat Asp	ata		-			gga				2901
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	Glu	aat			_	Asp	ata		-		Pro	gga				2901
Leu	Glu 915	aat Asn	Ala	Thr	Ser	Asp	ata Ile	Leu	Val	Ser	Pro 925	gga Gly	Asn	Glu	His	2901
Leu	Glu 915 atc	aat Asn gtg	Ala	Thr	Ser	Asp 920	ata Ile ttg	Leu tgt	Val	Ser	Pro 925 gaa	gga Gly agt	Asn	Glu	His tat	
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Leu aaa Lys 930	Glu 915 atc Ile	aat Asn gtg Val	Ala cct Pro	Thr gaa Glu	gct Ala	Asp 920 ttg	ata Ile ttg Leu	Leu tgt Cys	Val cat His	cag Gln 940	Pro 925 gaa Glu	gga Gly agt Ser	Asn tct Ser	Glu cac His	His tat Tyr 945	
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Leu aaa Lys 930	Glu 915 atc Ile	aat Asn gtg Val	Ala cct Pro	Thr gaa Glu ggt	gct Ala 935	Asp 920 ttg Leu	ata Ile ttg Leu	Leu tgt Cys	Val cat His	cag Gln 940	Pro 925 gaa Glu gtc	gga Gly agt Ser	Asn tct Ser	Glu cac His	tat Tyr 945	2949

ctg cat gat	gag tcc	ctc tcc	aca	agc	cct	cat	ggg	gct	ctc	cac	cag	3093
Leu His Asp	Glu Ser	Leu Se	Thr	Ser	Pro	His	Gly	Ala	Leu	His	Gln	
980			985					990				
cac gtg act	gat att	cag cti	: cat	att	ctt	aaa	aat	agg	atg	tat	gga	3141
His Val Thr	Asp Ile	Gln Let	His	Ile	Leu	Lys	Asn	Arg	Met	Tyr	Gly	
995		1000)				1005					
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Cys Asp Ser	Val Lys	Gln Pro	Arg	Asn	Ile	Glu	Glu	Pro	Ser	Asn	Ile	
1010		1015			1	1020					1025	
cct gta tca	gag aag	gat tct	aat	tct	cag	gtg	gat	tta	agc	att	cgg	3237
Pro Val Ser	Glu Lys	Asp Ser	Asn	Ser	Gln	Val	Asp	Leu	Ser	Ile	Arg	
	1030			1	1035					1040		
	1030			1	1035				:	1040		
gtt act gat		att ggt	. gaa			caa	gta	gat			ggt	3285
gtt act gat Val Thr Asp	gat gat			ata	att				gga	agc		3285
Val Thr Asp	gat gat		Glu	ata	att			Asp	gga	agc		3285
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Val Thr Asp	gat gat Asp Asp 1045 tcc aat	Ile Gly	Glu	ata Ile 1050 gga	att Ile agt	Gln	Val aga	Asp	gga Gly 1055 gca	agc Ser gat	Gly	
Val Thr Asp	gat gat Asp Asp 1045 tcc aat	Ile Gly	Glu	ata Ile 1050 gga	att Ile agt	Gln	Val aga Arg	Asp	gga Gly 1055 gca	agc Ser gat	Gly	
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yal Thr Asp gat aca tct Asp Thr Ser 1060 aat gaa ttt	gat gat Asp Asp 1045 tcc aat Ser Asn	gaa gaa Glu Glu	ata Ile 1065 gac Asp	ata Ile 1050 gga Gly	att Ile agt Ser	Gln aca Thr gat Asp	Val aga Arg	gat Asp .070	gga Gly 1055 gca Ala	agc Ser gat Asp	Gly gag Glu	3333
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Asp Asn Glu Asp Pro Gln Val Asn Ile Val Glu Glu Asp Pro Leu Asn

Glu Glu Ala Asp Ser Ile Ser Asn Glu Asp Ser Ala Thr Asn Ser Ser

1110 1115 1120

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Glu Phe Pro Ser Gly Ser Ser Ser Thr Ser Ser Thr Pro Leu Ser Ser 50 55 60

Pro Ile Val Asp Phe Tyr Phe Ser Pro Gly Pro Pro Ser Asn Ser Pro 65 70 75 80

Leu Ser Thr Pro Thr Lys Asp Phe Pro Gly Phe Pro Gly Ile Pro Lys

85 90 95

Ala Gly Thr His Val Leu Tyr Pro Ile Pro Glu Ser Ser Ser Asp Ser
100 105 110

Pro Leu Ala Ile Ser Gly Gly Glu Ser Ser Leu Leu Pro Thr Arg Pro 115 120 125

Thr Cys Leu Ser His Ala Leu Leu Pro Ser Asp His Ser Cys Thr His
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Glu Ser Leu Gly Phe Gln Ser Asp Asp Leu Pro Gln Phe Gln Tyr Phe 165 170 175

195 200 205

Ser Arg Asn Lys Glu Met Pro Ile Asp Gln Lys Ser Leu Asn Lys Cys
210 215 220

Ser Leu Asn Tyr Ile Cys Glu Lys Leu Glu His Leu Gln Ser Ala Glu 225 230 235 240

Asn Gln Asp Ser Leu Arg Ser Leu Ser Met His Cys Leu Cys Ala Glu 245 250 255

Glu Asn Ala Ser Ser Phe Val Pro His Thr Leu Phe Arg Ser Gln Pro
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Lys Ser Gly Trp Ser Phe Met Leu Arg Ile Pro Glu Lys Lys Asn Met
275 280 285

Met Ser Ser Arg Gln Trp Gly Pro Ile Phe Leu Lys Val Leu Pro Gly
290 295 300

Gly Ile Leu Gln Met Tyr Tyr Glu Gln Gly Leu Glu Lys Pro Phe Lys 305 310 315 320

Glu Ile Gln Leu Asp Pro Tyr Cys Arg Leu Ser Glu Pro Lys Val Glu 325 330 335

Asn Phe Ser Val Ala Gly Lys Ile His Thr Val Lys Ile Glu His Val $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$

Ser Tyr Thr Glu Lys Arg Lys Tyr His Ser Lys Thr Glu Val Val His

Asp Phe Leu Asp Phe Leu Thr Thr Val Glu Glu Leu Met Lys Leu Pro Ala Val Ser Lys Pro Lys Lys Asn Tyr Glu Glu Glu Glu Ile Ser Leu Glu Ile Val Asp Asn Phe Trp Gly Lys Val Thr Lys Glu Gly Lys Phe Val Glu Ser Ala Val Ile Thr Gln Ile Tyr Cys Leu Cys Phe Val Asn Gly Asn Leu Glu Cys Phe Leu Thr Leu Asn Asp Leu Glu Leu Pro Lys Arg Asp Glu Ser Tyr Tyr Glu Lys Asp Ser Glu Lys Lys Gly Ile Asp Ile Leu Asp Tyr His Phe His Lys Cys Val Asn Val Gln Glu Phe Glu Gln Ser Arg Ile Ile Lys Phe Val Pro Leu Asp Ala Cys Arg Phe Glu Leu Met Arg Phe Lys Thr Leu Tyr Asn Gly Asp Asn Leu Pro Phe Ser Leu Lys Ser Val Val Val Gln Gly Ala Tyr Val Glu Leu Gln

Ser Leu Arg Ser Cys Asp Asn Ile Arg Ile His Phe Pro Val Pro Ser 565 570 575

Gln Trp Ile Lys Ala Leu Trp Thr Met Asn Leu Gln Arg Gln Lys Ser 580 585 590

Leu Lys Ala Lys Met Asn Arg Arg Ala Cys Leu Gly Ser Leu Gln Glu
595 600 605

Leu Glu Ser Glu Pro Val Ile Gln Val Thr Val Gly Ser Ala Lys Tyr
610 620

Glu Ser Ala Tyr Gln Ala Val Val Trp Lys Ile Asp Arg Leu Pro Asp 625 630 635 640

Lys Asn Ser Ser Leu Asp His Pro His Cys Leu Ser Tyr Lys Leu Glu 645 650 655

Leu Gly Ser Asp Gln Glu Ile Pro Ser Asp Trp Tyr Pro Phe Ala Thr
660 665 670

Val Gln Phe Ser Val Pro Asp Thr Cys Ala Ser Arg Thr Glu Val Arg
675 680 685

Ser Leu Gly Val Glu Ser Asp Val Gln Pro Gln Lys His Val Gln Gln 690 695 700

Arg Ala Cys Tyr Asn Ile Gln Pro Lys Leu Tyr Arg Ser Val Ile Glu 705 710 715 720 740 745 750

Gln Ser Lys Ala Thr Glu Asp Phe Phe Arg Asn Ser Ile Gln Ser Pro
755 760 765

Leu Phe Thr Leu Gln Leu Pro His Ser Leu His Gln Thr Leu Gln Ser
770 780

Ser Thr Ala Ser Leu Val Ile Pro Ala Gly Arg Thr Leu Pro Ser Phe 785 790 795 800

Thr Thr Ala Glu Leu Gly Thr Ser Asn Ser Ser Ala Asn Phe Thr Phe
805 810 815

Pro Gly Tyr Pro Ile His Val Pro Ala Gly Val Thr Leu Gln Thr Val

Ser Gly His Leu Tyr Lys Val Asn Val Pro Ile Met Val Thr Glu Thr 835 840 845

Ser Gly Arg Ala Gly Ile Leu Gln His Pro Ile Gln Gln Val Phe Gln 850 855 860

Gln Leu Gly Gln Pro Ser Val Ile Gln Thr Ser Val Pro Gln Leu Asn 865 870 875 880

Pro Trp Ser Leu Gln Ala Thr Thr Glu Lys Ser Gln Arg Ile Glu Thr
885 890 895

Val Leu Gln Gln Pro Ala Ile Leu Pro Ser Gly Pro Val Asp Arg Lys

His Lys Ile Val Pro Glu Ala Leu Leu Cys His Gln Glu Ser Ser His Tyr Ile Ser Leu Pro Gly Val Val Phe Ser Pro Gln Val Ser Gln Thr Asn Ser Asp Val Glu Ser Val Leu Ser Gly Ser Ala Ser Met Ala Gln Asn Leu His Asp Glu Ser Leu Ser Thr Ser Pro His Gly Ala Leu His Gln His Val Thr Asp Ile Gln Leu His Ile Leu Lys Asn Arg Met Tyr Gly Cys Asp Ser Val Lys Gln Pro Arg Asn Ile Glu Glu Pro Ser Asn Ile Pro Val Ser Glu Lys Asp Ser Asn Ser Gln Val Asp Leu Ser Ile

Arg Val Thr Asp Asp Asp Ile Gly Glu Ile Ile Gln Val Asp Gly Ser

Gly Asp Thr Ser Ser Asn Glu Glu Ile Gly Ser Thr Arg Asp Ala Asp
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Glu Asn Glu Phe Leu Gly Asn Ile Asp Gly Gly Asp Leu Lys Val Pro \$1075\$

Ser Asp Asn Glu Asp Pro Gln Val Asn Ile Val Glu Glu Asp Pro Leu 105 1110 1115 1120

Asn Ser Gly Asp Asp Val Ser Glu Gln Asp Val Pro Asp Leu Phe Asp 1125 1130 1135

Thr Asp Asn Val Ile Val Cys Gln Tyr Asp Lys Ile His Arg Ser Lys
1140 1145 1150

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cagcatcaat aaaactgaca gecaaaaaaa a 261

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Ala Phe Pro Arg Arg Thr Ser Phe Asn Thr

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18

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